

FIGURE 9

1 GGATCCTATGTGAGGTTATAGATCCACGTGCATATCAAATTTCTCAAATTCGTTGCCTTC
-----+-----+-----+-----+-----+-----+-----+
CCTAGGATACACTCCAATATCTAGGTGCACGTATAGTTTAAAGAGTTTAAGCAACGGAAG
61 TAACGGCTCTTATATCCCTTCTCCGTGGACAGAATCTTCAGTCAAAAGCTCTGAGCTATG
-----+-----+-----+-----+-----+-----+-----+
ATTGCCGAGAATATAGGGAAGAGGCACCTGTCTTAGAAGTCAGTTTTCGAGACTCGATAC
N L Q S K A L S Y G
121 GTTCACACAATGAGTTACTTGAACCTGTGGATAGGTTAGTACTACCTGAGACTTTTTTCC
-----+-----+-----+-----+-----+-----+-----+
CAAGTGTTACTCAATGAACCTGAACACCTATCCAATCATGATGGACTCTGAAAAAAGG
S H N E L L E L V D
181 CTCTCCTCCTTTCATTACAAAGGTATTAGGGTTTCTTGTCAATCTGTGCATATATATGCA
-----+-----+-----+-----+-----+-----+-----+
GAGAGGAGGAAAGTAATGTTTCCATAATCCCAAAGAACAGTTAGACACGTATATATACGT
S
241 GCAAGCTTGTGGAATCAAATGTCCGGTGGTGAAGCGTGGACACCCTCGTTTCAGCTGGAGG
-----+-----+-----+-----+-----+-----+-----+
CGTTGGAACACCTTAGTTTACAGCCACCACATTTCGCACCTGTGGGAGCAAGTCGACCTCC
K L V E S N V G G V S V D T L V Q L E G
301 GTGTCCTTGAAAATGCCCTCTCTCTAACTAGAGCTAGGAAGGTACGTTGACTTCATACTG
-----+-----+-----+-----+-----+-----+-----+
CACAGGAACTTTACGGGAGAGAGATTGATCTCGATCCTTCCATGCAACTGAAGTATGAC
V L E N A L S L T R A R K
361 TCTTCTCATTTCTTACTTTGTTTGTGAAACGATTGTTCACTTATATTTAATTTGTTGCA
-----+-----+-----+-----+-----+-----+-----+
AGAAGAGTAAAGAATGAAACAAACAACTTTGCTAACAAGTGAATATAAAATTAACAACGT
421 GACAGAACTAATGTTGAAGCTTGTGATAGCCTCAAAGAAAAGGTTAGATATATCATATA
-----+-----+-----+-----+-----+-----+-----+
CTGTCTTGATTACAACCTTCGAACAACTATCGGAGTTTCTTTTCCAATCTATATAGTATAT
T E L M L K L V D S L K E K
481 TGATTTTATAGCACTTCAGATATCTTCTCGTGTGTTGAAAGCCTCAAATATTTATGTTG
-----+-----+-----+-----+-----+-----+-----+
ACTAAAATATCGTGAAGTCTATAGAAGAGCACAACTTTTCGGAGTTTATAAATACACAAC
541 TATTAAAGTTTCTCTAAGTGTGCTTTATGAGCTCGCAATCAAACCTTCTTCATAAGTGCATC
-----+-----+-----+-----+-----+-----+-----+
ATAATTCAAAGAGATTACACGAAATACTCGAGCGTTAGTTTGAAGAAGTATTCACGTAG
601 TGGTCTTTCANGGATGATTAAAAATATTGTTTTGGATACCAGAATCTGAAAATANGNTTA
-----+-----+-----+-----+-----+-----+-----+
ACCAGAAAGTNCCTACTAATTTTATAACAAAACCTATGGTCTTAGACTTTTATNCNAAT
661 AAAACTTGCAACTGATGAACATGTCCTTCAN
-----+-----+-----+-----+-----+-----+-----+
TTTTGAACGTTGACTACTTGTACAGGAAGTN 691

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